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Table 1. Cotransfections on 293 and 293Cre cells for rescue of LacZ (± loxP)

Plasmid combo	µg DNA	Plaques/dish (293 cells) (Totals	Plaques/dish (293Cre cells)	(Totals)
рСА36:pBHG10	5:5	0, 0, 0, 0	0, 1, 2, 0	
	5:10	0, 0, 0, 1	1, 0, 0, 0	
	10:10	2, 0, 1, 1	1, 2, 0, 0	
		(5)		(7)
pCA36:pBHG10lox	5:5	0, 0, 0, 1	0, 0, 0, 0	
	5:10	0, 0, 0, 1	0, 0, 0, 0	
	10:10	0, 0, 2, 1	0, 0, 0, 0	
		(5)		(0)
pCA36lox:pBHG10	5:5	1, 3, 1, 0	0, 1, 0, 1	
	5:10	0, 1, 0, 0	0, 0, 1, 2	
	10:10	0, 0, 0, 0	0, 1, 1, 0	
		(6)		(7)
pCA36lox:pBHG10lox	5:5	1, 0, 0, 1	15, 14, 20, 20	
	5:10	0, 0, 0, 0	11, 15, 12, 16	
	10:10	0, 0, 1, 1	18, 9, 10, 8	
		(4)		(168)

Table 2. Cotransfections on 293 and 293Cre cells for rescue of LacZ (\pm loxP)

Plasmid combo	μg DNA	Plaques/dish (293 cells) (Totals)	Plaques/dish (293Cre cells) (Totals)
pCA36:pBHG10lox	5:5	1, 1, 2, 6, 2, 3	1, 1, 2, 1, 2, 3
		(15)	(10)
pCA36lox:pBHG10lox	5:5	1, 2, 2, 2, 2, 1	41,44,41,41,44,31
		(10)	(242)
pCA36loxΔ:pBHG10lox	5:5	0, 0, 0, 0, 0, 0	41,36,55,34,24,40
		(0)	(230)
FG140	1	72, 72	150, 115

Table 3 Efficiency of Ad vector rescue by cotransfection with pBHG10lox and various shuttle plasmids

Cell line	Shuttle plasmid	Plaques/dish	Average/dish
293	pCA36lox	6, 2, 3, 3, 5	3.8
	pCA36loxA	1, 4, 0, 0, 0	1.0
	pCA36lox∆CreR	2, 2, 4, 3, 2	2.6
	pCA36loxACreT	9, 4, 4, 7, 3	5.4
293Сте	pCA36lox∆	23, 28, 22, 28	25.3

^{* 5}µg of all plasmids were used in cotransfections.

Table 4. Efficiency of Ad vector rescue by cotransfection of 293 cells with pBHG10lox and shuttle plasmids encoding Cre*.

Cell line	Shuttle plasmid	Plaques/dish	Average/dish
293	pCA36lox	2, 3, 1, 0, 1	1.4
	рСА361охА	1, 0, 0, 0, 0	0.2
	pCA36lox∆CreT°	3, 1, 5, 2, 4	3.0
	pCA36loxCreFTRb	21, 20, 42, 34, 40	31.4

 $^{^{3}}$ All corransfections performed with 5 μg of the indicated shuttle plasmid and 5 μg of pBHG10lox

^bPlasmids illustrated in figure 8c.

Table 5. Efficiency of rescue of fibre and E4 genes into Ad by cotransfection with pFG173lox and pFG23lox^a

		Number of plaques (average/dish)	
Plasmids	μg DNA	293 cells	293Cre cells
pFG173lox ^h : pFG23dX1loxe ^c	5:5	0, 0, 0, 0 (0)	33, 27, 39, 26 (31)
•	2:2	0, 0, 0, 0 (0)	9, 15, 10, 9 (11)
pFG173: pFG23dX1	5:5	0, 0, 0, 0 (0)	0, 0, 1 (0.3)
pFG140	1	95	93

^{*}Cotransfections as diagrammed in figure 9

^bDiagrammed in figure 9b

Diagrammed in figure 10